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## **CLAIMS**

## What is claimed is:

- 1. An isolated polynucleotide comprising:
  - (a) a nucleotide sequence encoding a polypeptide having 1-FFT activity, wherein the amino acid sequence of the polypeptide and the amino acid sequence of SEQ ID NO:2, 4, or 6 have at least 90% sequence identity based on the Clustal alignment method, or
    - (b) the complement of the nucleotide sequence, wherein the complement and the nucleotide sequence contain the same number of nucleotides and are 100% complementary.
- 2. The polynucleotide of Claim 1 wherein the amino acid sequence of the polypeptide and the amino acid sequence of SEQ ID NO:2, 4, or 6 have at least 95% sequence identity based on the Clustal alignment method.
- 3. The polynucleotide of Claim 1 wherein the polypeptide comprises the amino acid sequence of SEQ ID NO:2, 4, or 6.
- 4. The polynucleotide of Claim 1 wherein the nucleotide sequence comprises the nucleotide sequence of SEQ ID NO:1, 3, or 5.
  - 5. A vector comprising the polynucleotide of Claim 1.
- 6. A recombinant DNA construct comprising the polynucleotide of Claim 1 operably linked to a regulatory sequence.
- 7. A method for transforming a cell comprising transforming a cell with the polynucleotide of Claim 1.
  - 8. A cell comprising the recombinant DNA construct of Claim 6.
- 9. A method for producing a plant comprising transforming a plant cell with the polynucleotide of Claim 1 and regenerating a plant from the transformed plant cell.
  - 10. A plant comprising the recombinant DNA construct of Claim 6.
  - 11. A seed comprising the recombinant DNA construct of Claim 6.
- 12. An isolated polynucleotide comprising a first nucleotide sequence,
  wherein the first nucleotide sequence contains at least 30 nucleotides, and wherein
  the first nucleotide sequence is comprised by another polynucleotide, wherein the
  other polynucleotide includes:
  - (a) a second nucleotide sequence, wherein the second nucleotide sequence encodes a polypeptide having 1-FFT activity, wherein the amino acid sequence of the polypeptide and the amino acid sequence of SEQ ID NO:2, 4, or 6 have at least 90% sequence identity based on the Clustal alignment method, or

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- (b) the complement of the second nucleotide sequence, wherein the complement and the second nucleotide sequence contain the same number of nucleotides and are 100% complementary.
- 13. An isolated polypeptide having 1-FFT activity, wherein the amino acid sequence of the polypeptide and the amino acid sequence of SEQ ID NO:2, 4, or 6 have at least 90% sequence identity based on the Clustal alignment method.
- 14. The polypeptide of Claim 13, wherein the amino acid sequence of the polypeptide and the amino acid sequence of SEQ ID NO:2, 4, or 6 have at least 95% sequence identity based on the Clustal alignment method.
- 15. The polypeptide of Claim 13, wherein the amino acid sequence of the polypeptide comprises the amino acid sequence of SEQ ID NO:2, 4, or 6.
- 16. A method for isolating a polypeptide encoded by the polynucleotide of Claim 1 comprising isolating the polypeptide from a cell containing a recombinant DNA construct comprising the polynucleotide operably linked to a regulatory sequence.
  - 17. An isolated polynucleotide comprising:
    - (a) a nucleotide sequence encoding a polypeptide having 1-SST activity, wherein the amino acid sequence of the polypeptide and the amino acid sequence of SEQ ID NO:14 or 16 have at least 97% sequence identity based on the Clustal alignment method, or
    - (b) the complement of the nucleotide sequence, wherein the complement and the nucleotide sequence contain the same number of nucleotides and are 100% complementary.
- 18. The polynucleotide of Claim 17 wherein the polypeptide comprises the amino acid sequence of SEQ ID NO:14 or 16.
- 19. The polynucleotide of Claim 17 wherein the nucleotide sequence comprises the nucleotide sequence of SEQ ID NO:13 or 15.
  - 20. A vector comprising the polynucleotide of Claim 17.
- 21. A recombinant DNA construct comprising the polynucleotide of Claim 17 operably linked to a regulatory sequence.
- 22. A method for transforming a cell comprising transforming a cell with the polynucleotide of Claim 17.
  - 23. A cell comprising the recombinant DNA construct of Claim 21.
- 24. A method for producing a plant comprising transforming a plant cell with the polynucleotide of Claim 17 and regenerating a plant from the transformed plant cell.
  - 25. A plant comprising the recombinant DNA construct of Claim 21.

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- 26. A seed comprising the recombinant DNA construct of Claim 21.
- 27. An isolated polynucleotide comprising a first nucleotide sequence, wherein the first nucleotide sequence contains at least 30 nucleotides, and wherein the first nucleotide sequence is comprised by another polynucleotide, wherein the other polynucleotide includes:
  - (a) a second nucleotide sequence, wherein the second nucleotide sequence encodes a polypeptide having 1-SST activity, wherein the amino acid sequence of the polypeptide and the amino acid sequence of SEQ ID NO:14 or 16 have at least 97% sequence identity based on the Clustal alignment method, or
  - (b) the complement of the second nucleotide sequence, wherein the complement and the second nucleotide sequence contain the same number of nucleotides and are 100% complementary.
- 28. An isolated polypeptide having 1-SST activity, wherein the amino acid sequence of the polypeptide and the amino acid sequence of SEQ ID NO:14 or 16 have at least 97% sequence identity based on the Clustal alignment method.
- 29. The polypeptide of Claim 28, wherein the amino acid sequence of the polypeptide comprises the amino acid sequence of SEQ ID NO:14 or 16.
- 30. A method for isolating a polypeptide encoded by the polynucleotide of Claim 17 comprising isolating the polypeptide from a cell containing a recombinant DNA construct comprising the polynucleotide operably linked to a regulatory sequence.